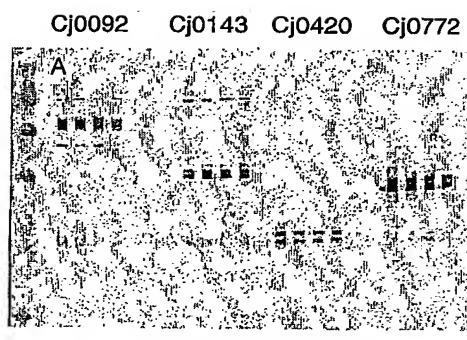
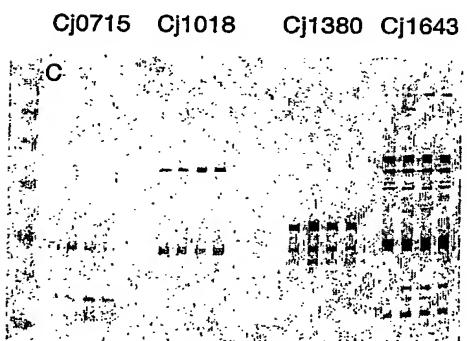


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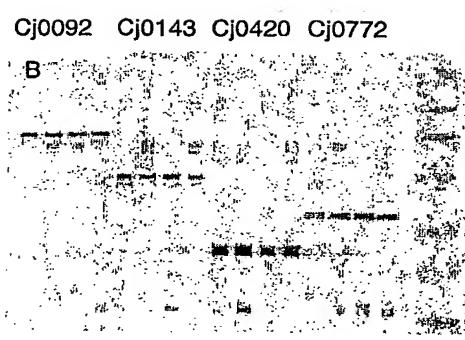
Fig. 1



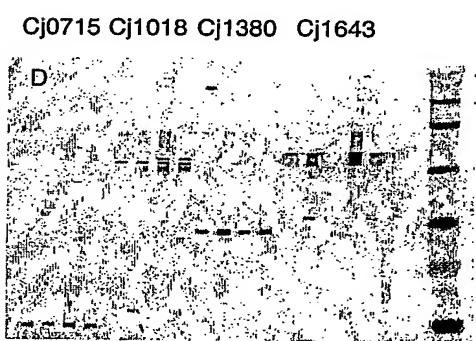
Recombinant proteins



Recombinant proteins



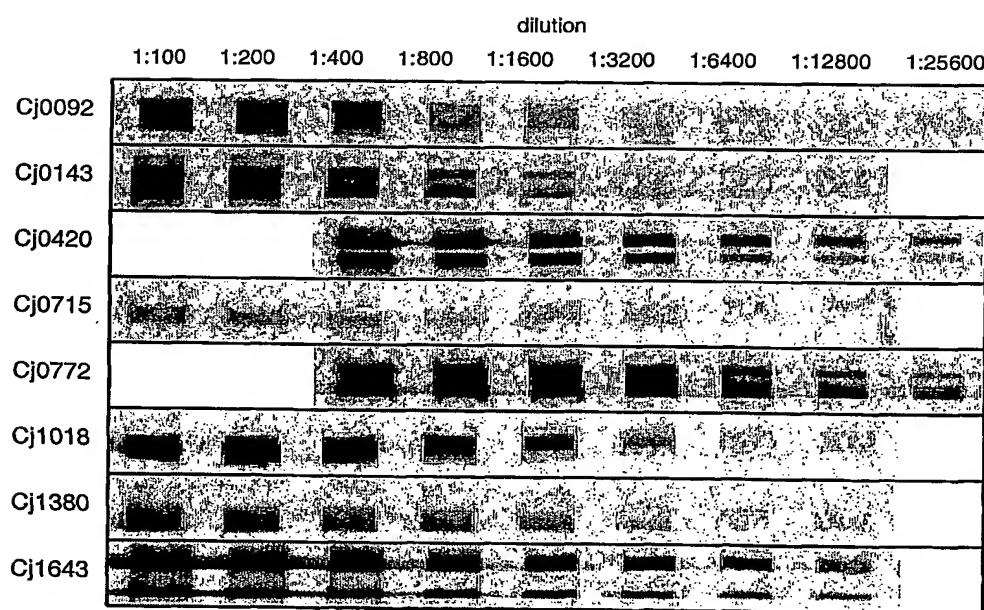
C. jejuni whole cell lysate



C. jejuni whole cell lysate

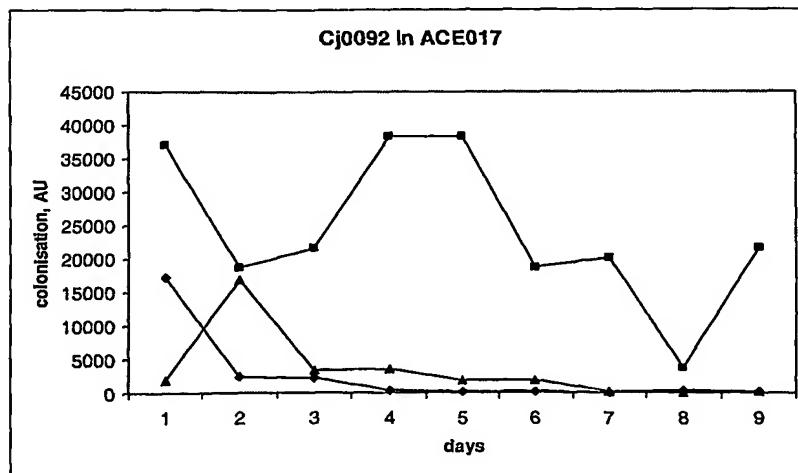
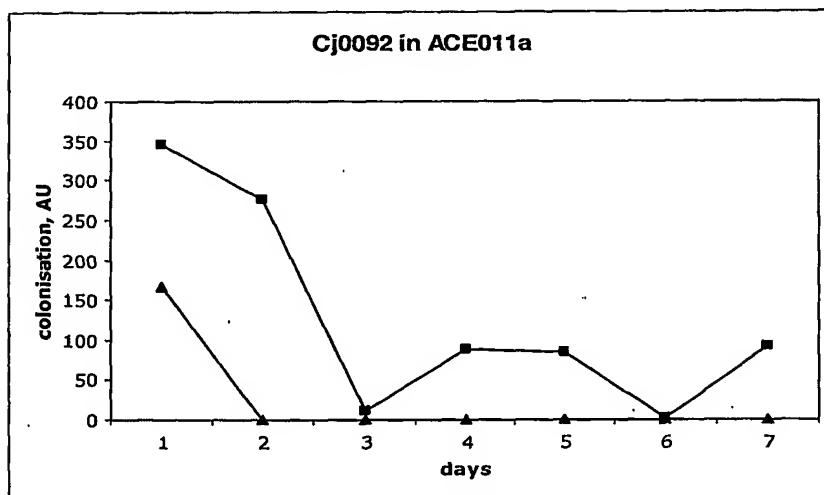
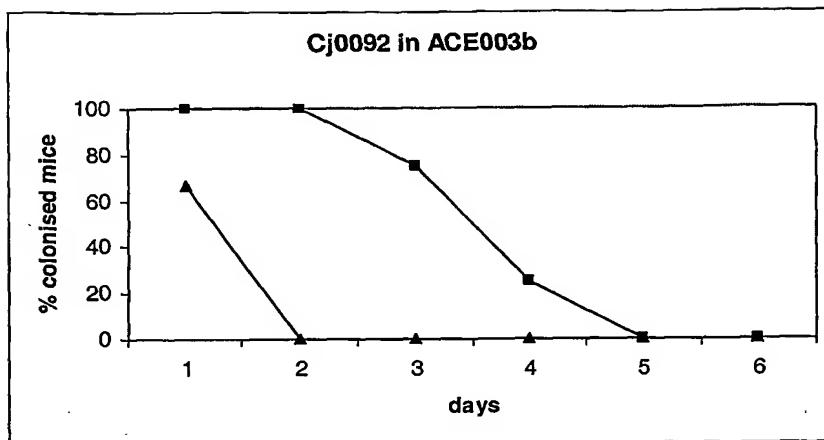
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Fig. 2



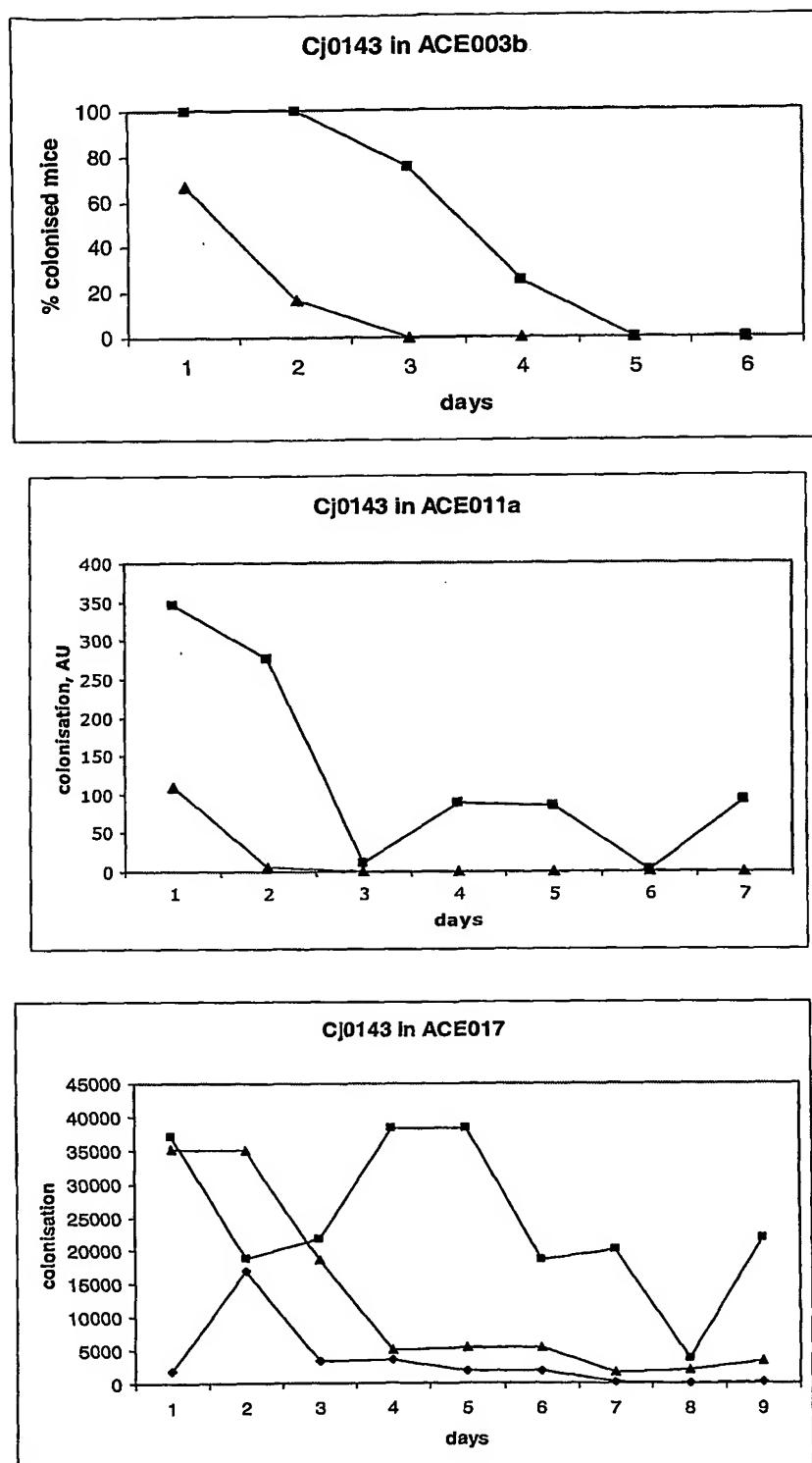
3/20

Fig. 3

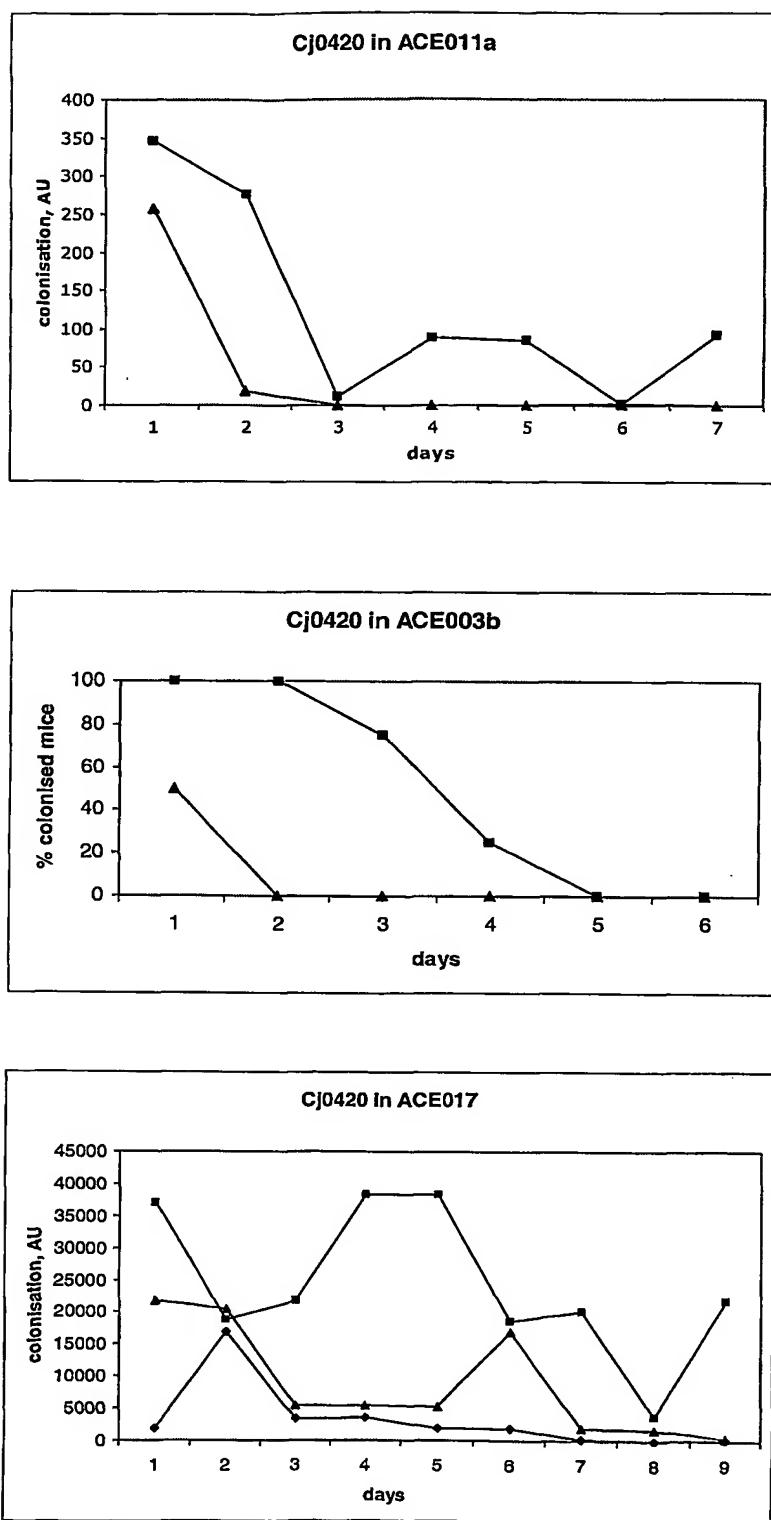


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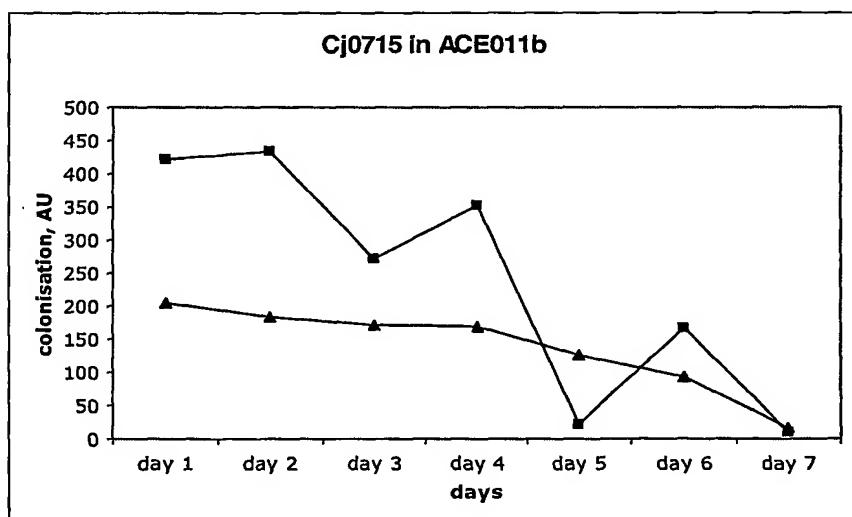
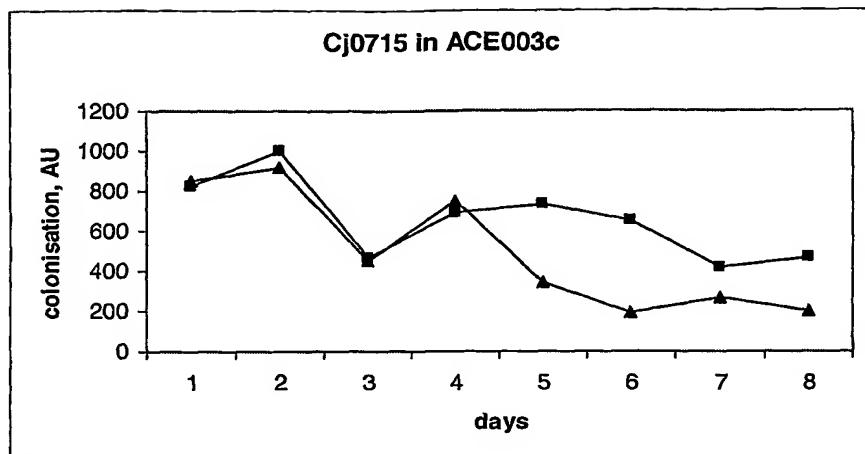
Fig. 3



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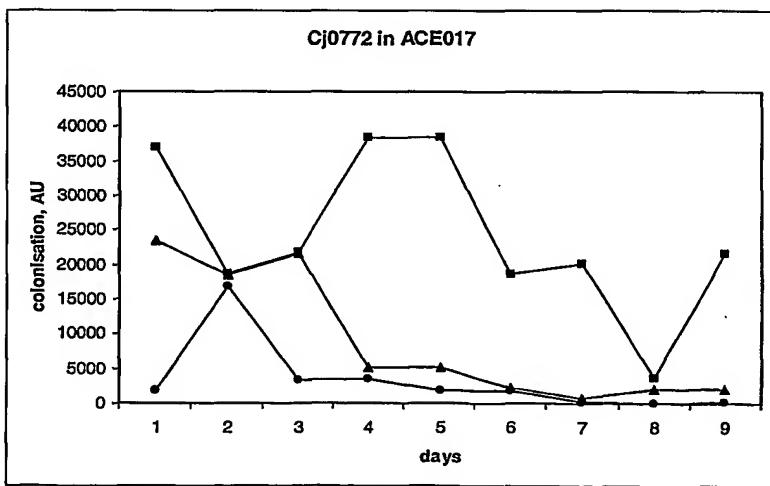
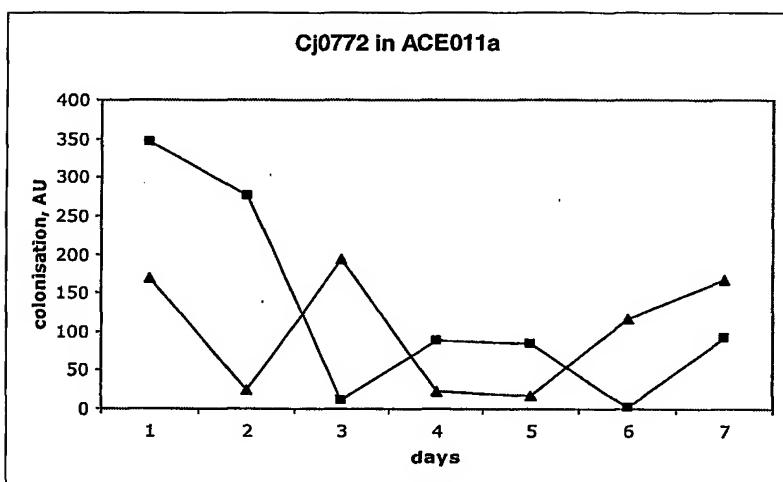
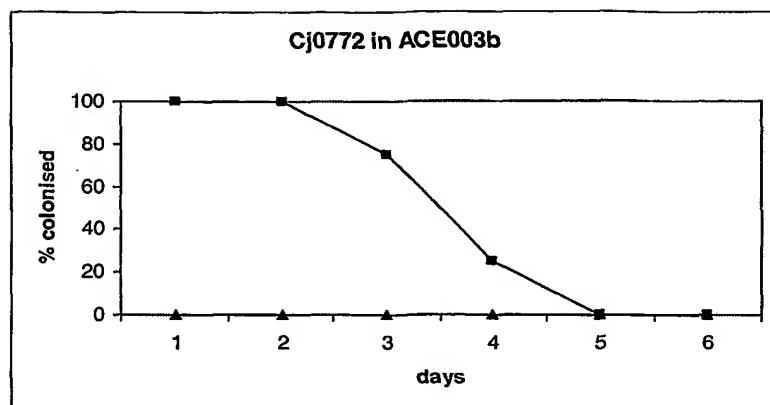
Fig. 3

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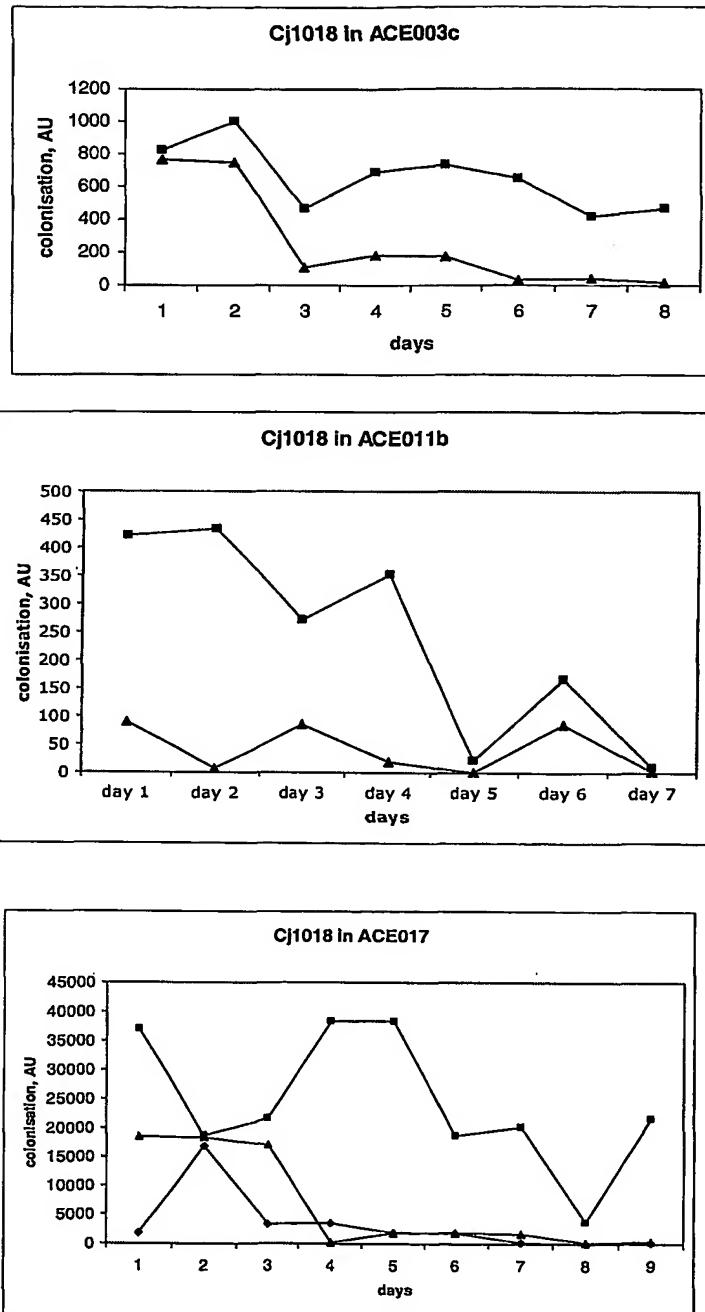
Fig. 3

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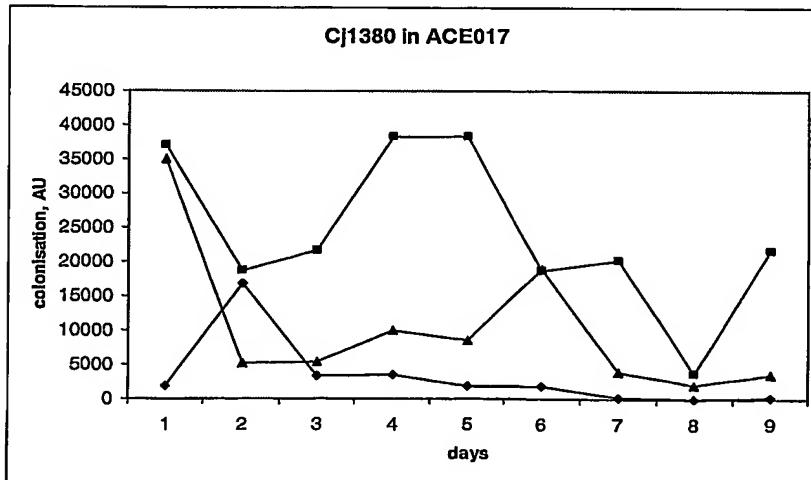
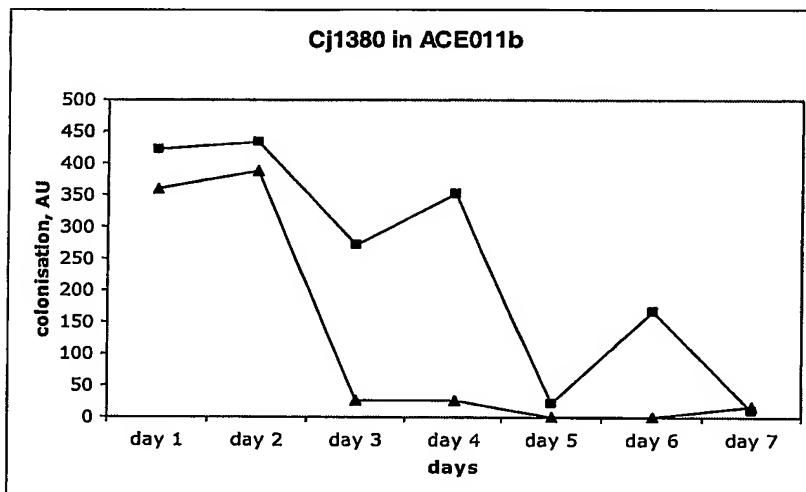
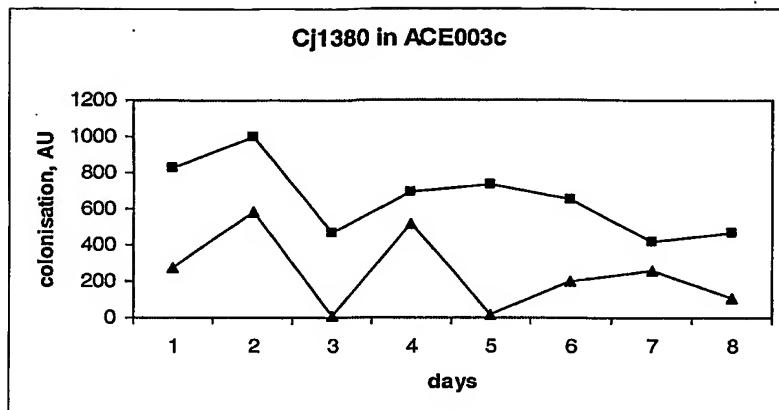
Fig. 3



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Fig. 3

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Fig. 3

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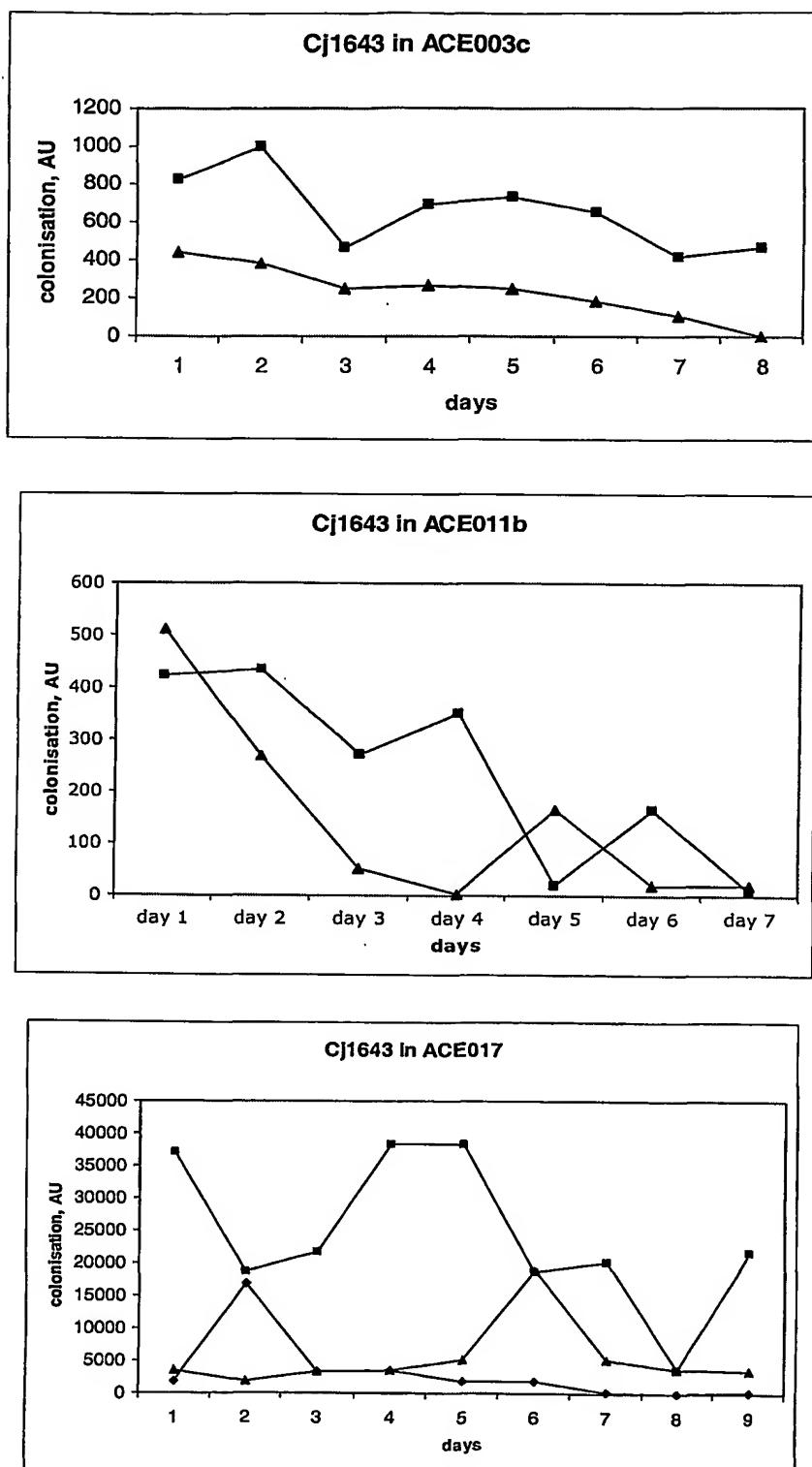
Fig. 3

FIG. 4

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Sequence listing

SEQ ID NO:1

ACE83 Cj0092

>AnrP630851 (NC_002163) putative periplasmic protein [C. jejuni]
 MKKILFIGSLVMASLLYAOGSQPVEITQQDINTQNEMSDASTKDITPKSIEDFFEEFADNFGIEYGITKD
 GKTFYTGKSTVAVNDTDPQFAQLQNAYQKAMLNQSEFIRDAFGRIATSKIQNYEADNSTNAKEFDELP
 KGDKDQILNKLTQLAGAQLDKALKDGLIDTNLSEDRKKTLLKQEFLNKMTNAIGSMSGLVPVQTIVT
 QRRGEYDVGVVAVISNKTRQLAKDMALARQSAIKGKGKAISEYLPKDTKGFLNEYGIRLVYDENGAPIIL
 SYGNWGVYADPSNAKKTNIILEDRAKETALTMADAIIIEFINTNLSQLKDERTTGDTYEEIIKQSINVNDSS
 TQEQTQNITNIIDKVNSKIKASASGKIRGIRTLKKWSYTSENGIEHVGAVRFYSYENLANTNEALNSKN
 ATKNEAKKSSSIQRSSNVVNSMDDF

SEQ ID NO:2

ACE 5 Cj0005

>AnrP544820 (NC_002163) putative molybdenum containing oxidoreductase [C.
 MKQNDQKENRRDFLKNIGLGLFGISVLSNFSENFLGSKALAKELPDFKIEGKLDIYHGKPLTAETEI
 YALDSDFTKPENFFVRNNNGLPPSLETIKERLHKGWTLEIDGESIINKSYTIEDLKKKFKTYTYALTVEC
 GGNRSEVIPSTKGTQWGYGAVACGRWTGVRKLDILKDCGIKNDAVYIGYYGIDTKLNGETSPISRGPV
 ISKALQDETTLIAWAYEGKDIPLVNGYPLRLVCGGYPASTSGKWLYKISVRNKHIDGEKMEGSYKVPVNPV
 KPGDFNYKGEMKIIESMPIRSVITNIKNGSEIKANKFEVRGKAWAGELEVSEVVSVNDYGVWTAKAVE
 KPLNRLAWQKWSAQISIPTKGYYEIWARAIDSQGNSQPMVLAQWNPGGYINNACHRVNVYGV

SEQ ID NO:3

ACE 29 Cj0029

>AnrP372217 (NC_002163) cytoplasmic L-asparaginase [C. jejuni]
 MKKAKSRIAIALGTGGTIAGFIDSTIATTGYAAGAIDIDVLIKAVPQIRDLADISWEQIANIDSSNCDEI
 WLRLAKKIAKLFAEGIDGVVITHGTDMEFTAYFLNLTIKSDFKPVVLVGAMRPTAISADGPKNLNAV
 LVVNKEAKNKGVMVAINDKILSARGVVKTHSLNDAFSSPDFGDLGYIVDGKVFYNNVIKAHTKNAPFD
 VSKLTSLEPKVDILYSYSNDGSGVAAKALFEHGTGIVVAGSGAGSIHKNQKDVLKELLKKGLKVVVSSRV
 VAGCVAVSDSDEKLGFISAEDLNQPKARVLLMLALTKTSDPKKIQEYFLKY

SEQ ID NO:4

ACE 37 Cj0037

>AnrP501075 (NC_002163) putative cytochrome c [C. jejuni]
 MKKHILLGLCLSLSLSAKSVDYKVGEELSKEGVEYFKELSKRPVQEWPKNLSSINDVPKGQGDLIR
 YGIELLSKTESTLGLPYSKLKKTSNEVNCISCHMDNDGNGLPGTKVYIPFLNILNNYPRLDIETMKIISV
 EDRIRGMGGTDSHRFPNDSKEMKAILAYFKWLKEAYGIKDGVKLEGDFFAKMNFPNRPADPVRGKKLFEE
 NCVACHGERGLGVKNDNYEQGSGHLYPSLLIYPDGGHMAMIPFLARFLKSAMPFGASADNPILSDEDALD
 IAAYVNTGFVRMPITTENRAGLDTAYSKSPSLKPEYFASPQQNLDPKEYIKVKYGPWKNPNHFPGF

SEQ ID NO:5

ACE 84 Cj0093

>AnrP666574 (NC_002163) putative periplasmic protein [C. jejuni]
 MKIIKILFLGLFLSLSNAKVITTTSTSSTGEGTGLTREDAINNAIEAIGKMSGVSINSLKKSNNTSV
 TDNSGSNIQDNYSEQISKATKGRADTYEINSVEQDANGKYTANVTIFKTTTKKYQAPGLSADNRRSITV
 FDSTPDAAKRGIGSAIQQKIISSLQSRKFNVLDRDSSGYYEMEKALIKSGDAASDEVYKLKNMLATDYI
 LLFSISGLEGKQKTSNLTKSKTEIEVIVDVRVLLFATRQIKFSNTLSMKVNLKDNLSANETALKQIAN
 RIAGDILNATIYPLKVASVENNEVIFSQSLSNQGDVYECFALGKVIKDTYTKENTGRVESKTGSIEITRTSP
 KFSYAKITEGGSVKVGDIICRPLSNTGSGNGYTIGRDANYQTQEGGGVNLGF

SEQ ID NO:6

ACE 98 Cj0107

>AnrP732169 (NC_002163) ATP synthase F1 sector beta subunit [C. jejuni]
 MQGFISQVLGPVVDVFNDYLPQINEAIVNFESEGKKHKLVLEVAHLGDNVRTIAMDMTDGLVRGLK
 AEALGAPISVPVGEKVLGRIFNVTGDLIDECEEISFDKKWAIHRDPPAFEDQSTKSEIFETGIKVVDLLA
 PYAKGGKVGLFGGAGVGKTVIIMELIHNVAFKHSGYVSFAGVGERTREGNDLYNEMKESNVLDKVALCYG
 OMNEPPGARNRIALTGLTMAEYFRDEMGLDVLMPIDNIFRFSQSGSEMSALLGRIPSAGVYQPTLASEMG
 KFQERITSTKKGSITSVQAVVVPADDLTDPAPATVFAHLDATTVLNRAIAEKGIYPAVDPLDSTSRLD
 NIIGEEHYKVARGVQSVLQKYKDLQDIIAIILGMDELSEEDKLVVERARKIEKFLSQPFFVAEVFTGSPGK
 YISLEDTIAGFKGILEGKYDHLPENAFYMVGNIDEAIAKADKLKG

FIG. 4

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SEQ ID NO:7

ACE 103 Cj0112

>AnrP511634 (NC_002163) periplasmic protein [C. jejuni]
 MKKIVAIIFLVFLGLSLWAEDPVIDVVNSGVVLPKIIVKDNLNSLSDENLKKSFYNNIVNDLKVSNNFEVVAN
 ATETSNYIIFYEYTLNKGNTLSNVKIKAGGSDKSEQTYTLNGLEQYPFLAHKSVKASVNALGLAPVDWMD
 HKILIARNSSSKKSQIIIMADYTLTYQKVIVDGLNLFPKGWNKEQTLFYYTAYDHDKPTLYRYDLNTNKA
 SKILSSGGMVVASDVNVDGSKLLVTMAPKDQPDVLYDLNTKNLTQLTNYSGIDVNGNFIGSDDSKVVFV
 SDRLGYPNITFMQDLNSNSAEQVFHGRNNSAVSTYKDFLVYSSREPNQAGVFNIYLMINSIDYIRQLTAN
 GKNLFPRFSSDGGSIVFIKYLGAAQSALGVIRVNANKTFYFPLRVGKIQSIDW

SEQ ID NO:8

ACE 134 Cj0143

>AnrP57234 (NC_002163) periplasmic solute binding protein for ABC
 transport system [C. jejuni]
 MKKIIILFILSLGIFYFTQAKNLEQEQTSSNLVSVSIAPIQAFFVKKIAANTLDVNVLPPNSNEHNFEF
 KPSTMKKLEKSDIYFTIGLEFEKVFTDKFQKQFPKLQVINMQKNIALIQTHDTHEHSHEHHHEGHFDP
 HTWLDPILVQTMALNITYDTLIQKYPQNENLYKENLDKFLAELDSLNQIASKLEKLKNREFVYHPSWTY
 FAKRYNLTQIPVEILGKEPKSKDLQKLITLMKDKNLKVIIFVQNGFPENAATLAKECDAKIYKIDHLSYD
 WENELLKTADAFSHNL

SEQ ID NO:9

ACE 159 Cj0169

>AnrP829849 (NC_002163) superoxide dismutase (Fe) [C. jejuni]
 MFELRKLPYDTNAFGDFLSAETFSYHHGKHNTYVTNLNNLIKDTFAGKDLVSIIKTSNGGVFNNAQV
 YNHDFYFDCIKPSTGCGCGGSCSIDANLQAALEKEGSLENFKAEFIKGATGVFGSGFWLVYNTKNQK
 LEFGVGTNSAATPITEDKVPLLVDVWEHAYVDRNARPAYLEKFYAHINWEFVAKAYEWALKEGMGSVS
 FYANELHPVK

SEQ ID NO:10

ACE 183 Cj0193c

>AnrP139712 (NC_002163) trigger factor (peptidyl-prolyl cis /trans
 isomerase, chaperone) [C. jejuni]
 MEVKAKQLDSVNATAVKIPSGMIKSEVENLAKKASKSVKMDGFRPGKVPVSAVLKRYERELTQDAEQNL
 FKSAVNSALQELKKENKELVGEPYFEKFDRKDGEIIAELILSFKPEIKLEGYEKLIPEYQTPKVSKKEID
 EKKDELLKRFATPEAIKTKRALKEGDFAKFDSEGFDKAFEGGKAENYVLEIGSKQFIPGFEDGMVGMK
 IGEEKDIKVTFPKEYGAHLLAGKDAVFVKVLHEIQELKIPELDDEMCKLLPGEEKASVEVLDEKLKEQI
 KNEKLFKLVNDDELKGKFADALIEKYNFDLPKGIVEQETDMQMRAAFNTFSEKEIEELKASKEKYQEKRDS
 FKEEAQKSVKLTIFIIDELAKLRKIEVNDQELIQAIYFEARYGMNPKEHENYKKQGALPAVKMALIEEK
 LFNDIFIPKTEKSEKVKKEKEDK

SEQ ID NO:11

ACE 259 Cj0285c

>AnrP467527 (NC_002163) chemotaxis protein [C. jejuni]
 MFDENIVKTSNELEMVLDFRIFKQGHDKVYEGIYGVNVSKVREIIKIPSLTELPGVPDYIEGIFDLRGVV
 IPVVNLAKWMQITEPESTMLKPRVIITEFSNLIIGFIVHEAKRIRRINWKDIEPATFSTGSGALDKGKIT
 GVTRIENDEVLLILLDLESVVEDILGIYAPKTDIDFGKIEKFTGTALILDDSMTARKRVKEMMQMGFQVVE
 AKDGVEGINKLEELSQIYGESLNDTLKIIIVSDVEMPQMDGFHFAARIKEDPRFKDIPIVFNSSLNEFMN
 EKGVQEAGGEGYLVKFNASDFFNEIAKVIKKHQSEQG

SEQ ID NO:12

ACE 332 Cj0358

>AnrP681041 (NC_002163) putative cytochrome C551 peroxidase [C. jejuni]
 MKVKSLLIASLVAFFSSLNAASLIDEAKNSGLVALPKDQKGVDIELKQNGVKASEFTLEKAELGKKLYFEP
 RLSKSGIISCNCTCHNVGLGGTDGISTAIGHKWTANPHHLNSPTVYNAVLNNTQFWDGRAGTLADQAKGPI
 QADPEMATPAKLAKEVKISSLPEYVSEFKKIYKGSGVNFNDIADAIAFERTLITPSRFDKFLEGDEKALT
 KEEQKGLKLFIDKGCVACHNGVNLGGNMQAFEVAGKYKFANLGDFKGDANGMVKPTLRNAETAPYFHN
 GAIWNLKDAIKEMGSVQLGIKISDKEAKSIEFLKSLTGKPAIVYPQLPISTEKTPKPEL

SEQ ID NO:13

ACE 419 Cj0448c

>AnrP569688 (NC_002163) putative MCP-type signal transduction protein
 [C. jejuni]
 MFGSKINHSDLQKLEENKNLTHKIEKFQSENLELKNKITSLEQAALESKLKTDLLNVLLTGVLKNNITII
 QGDMLENVNKAEVISYSKTSLAEMDELNHANSINASLGNITESANKTRDVAGTLHRSVDEITNVINLI
 KDVSQDTNLLALNAAIEAARAGEHGRGFAVADEVRLAEGTQKATTEVEMNINLLKQANEMYTQSEQV
 EKISIDSNAHIMSFSKEKFTHLVNEAHSTNSAVGIASEAFVSLAKLDHIAFKLNGYKEIFSKSGKQLADH
 TSCRRLGKWLASTGKERFGQNKSLKINEPHEKVHENMNNAITIANTEDISKDITQHSIINKCEVAENASL

FIG. 4**13/20**

DLFNVFKEMLDESDH

SEQ ID NO:14

ACE 420 Cj0449c

>AnrP852550 (NC_002163) hypothetical protein Cj0449c [C. jejuni]
MLHEYRELMSELKGKDAHFDKLFDRHNELDDMIKDAEEGRTSLSSMEISTLKKEKLHVVKDELSQYLANYK
K

SEQ ID NO:15

ACE 481 Cj0511

>AnrP255677 (NC_002163) putative secreted protease [C. jejuni]
MMELILKTKRFFAGLAGFATTFLCLFLTSHLQAKVDQKEEQVQKRLEALDKLTKTLLAIVEOYYVDDONI
SDLVDKSLSGGLSNLDAHSSFLNEKDFNDMKIQTNGEFGGLGITVGMKDGAATVVSPIEGTPADKAGIKS
GDIILKINDEATLGGINLNDAVDKMRGKPKTQITLTIFRKGATKPFVTLTREIIKIESVYAKMIENENIL
YLRVTVNFKNVVVDVASKELKKYPNVKGVILDRLRNPNPGGLLNQAIGLVNLFVDKGVIVSQKGRIASENQEY
KADPKNKISNASLVLVNGGSASASEIVSGALQDLKRGVIVGENTFGKGSVQQIIPINKTEALRLTIARY
YLPMSGRTIQAVGVKP DIEVFPGVNTQEDGFSIKESDLKQHLESELEKIDKNKEDKQENKDNKNLISQK
QINDDAQLKSAIDTIKILNIKQGQ

SEQ ID NO:16

ACE 528 Cj0559

>AnrP252410 (NC_002163) oxidoreductase [C. jejuni]
MKKIDLIVVGAGPTGIGCAVEAKLKNKEVLLILEKSNNICQTLMQFYKDGKRVDKAYKGCEGTNHGHVPFE
DGTKESTIETFQNALKHEHNEVEFGSEVESVKNENGVFLVSTAKGVYECKNIIVAIGRMGKPNKPDYKLP
MTLTKIIMFNANSVLGNENKEKILVGGGNSAAEYAVDLANSNQVSLCYRKKEFTRLNDINLKDHEAGNSK
VELKLGIDINEVEDDNGAKVNFTDGTSDIYDRIIYAIGGSTPLDFLQKCGINVDDKGVPPLMDENKQSNV
KGIFVAGDIATKNGASIVTGLNDAVKILSVL

SEQ ID NO:17

ACE 582 Cj0613

>AnrP916533 (NC_002163) possible periplasmic phosphate binding protein
[C. jejuni]
MKKILSLSVTSLALCGALNAVDLKIAGSSTVYPFTSFVAEEYASIKNTKTPIVESLGTGGGFKVFC EGT
DISNASRPMKLSEFETCKKAGVT DIVGMMIGYDGVLAQNKTNAPLNITKKELFLALAKEIPONGKLI P
PYTNWNQINKNLPNRKISVYGPSSSGTRDTIEELVMSDVSKKIPEYKGEYKTRQDGAYIPSGENDNLI
VSKLTIDKAFIGIFGYSFVNSDKINAANIDGVTPESESIADEKYELARSLFIYINAKKNPKEAFDFAK
IYMSDDLAKSGGELEKIGLVPLOSSDKLAKASQKHVEDRKILDELVKAGKVF

SEQ ID NO:18

ACE 605 Cj0636

>AnrP126795 (NC_002163) NOL1\NOP2\sun family protein [C. jejuni]
MQNILSSFAQEKNVCVFANTLKTSIEELEKEFLQNLKFKKINVYCYLFDAKDKA ILSSMKAFNEAHFYI
QNYSSYLCALNLEVKGQS VLDMAAAPGGKSINLANFMQNTGYLACNEMSRDRFFILQKNLKNYGVNAKV
FMKDGNIGNLCPFKDILL DAPCSTFAKIGFDLEKSYKEIKNIAKTQKKLLHSALKALKIGGELVYST
CTFTKEENEVIEVIAKLSEFKLELL DIDLENVEAKAGQSEEF AEISKRRILPSLDYDGFFIAKLRKLC

SEQ ID NO:19

ACE 667 Cj0706

>AnrP327756 (NC_002163) hypothetical protein Cj0706 [C. jejuni]
MNKYLEQLVLLSKIDQEIDSYEPKIDSINKTLKDAELKIEKINADLEKIDEIKDIENQKIQNNAHISEP
SAKIKDL SKKSGVVKTEKEANALKIEEDI AKEQLDAANDEIVRLDKILEN KETYKKELEEEKIKQEONIN
EIRVSIKSEM EVLEKDRMSVYDKTKLVSE MNQKVLSFYEKIRKWA NTAVV PVKKQACYGCFM KIYDKT
YLSVVKGEEIVTCPHCGRILYKEQEEQN

SEQ ID NO:20

ACE 676 Cj0715

>AnrP684299 (NC_002163) transthyretin-like periplasmic protein
[C. jejuni]MFSIKKTLLILASVPMFLSATEYQLSTHVLDITSGQPAPKVKVELYKLEANQQWKKVSEEFTEENGRIGD
LLPYEKAENRAFGIYKLKFFT KDY TSHKINTFYPFVEVS FELSKDQHYHVPITLSPFGYSTYRGS

SEQ ID NO:21

ACE 731 Cj0779

>AnrP191193 (NC_002163) probable thiol peroxidase [C. jejuni]

MSIVNFKGPNVKLGNSVEVGADAPKVN LKAKDLSVIEIGAAGKTQJILSVP SL DTPVCATEAREFNKKV
ASYNGAEVIVVSM DLPFAMGRFCSTEGIENLSVASDFVAKEFGEKYGVLINEGALEGLLARAVFVIKEGK
VAYKELVNEITEMPDIAKLDAFFGGSSCCGGCGCH

FIG. 4**14/20**

SEQ ID NO:22

ACE 853 Cj0909

>AnrP318705 (NC_002163) putative periplasmic protein [C. jejuni]
 MKKILLLGALFAVNWLAVNDIEVKNAFVKQTPPPHAQNSAIFLTIFNNTNKDIALISAKSDISEVSELHTH
 IHKGKMMMQKIPPEIIIAKSSTELKSGGYHIMLLKLKKPIIKDTKVNLSDLKFNNHKTIELKNIDSKEF

SEQ ID NO:23

ACE 937 Cj0994c

>AnrP493933 (NC_002163) ornithine carbamoyltransferase [C. jejuni]
 MKHFLTLRDFSKEEILSLVNHASELKKEPKLQLQDKTLAMIFEKNSTRTRMAFELAITELEGGKALFLSSN
 DLQLSRGEVKDTARVIGAMVDFVMMRVNKHETLLEFARYSKAPVINALSELYHPTQVLGDLFTIKEWNK
 MQNGIAKVAFIGDSNNMCNSWLITAAILGFEEISIAMPKNYKISPEIWEFAMKQALISGAKISLGYDKFEA
 LKDKDVVITDTWVSMGEENEKERKIKEFEGFMIDEKAMSVANKDAILLHCLPAYRGYEVSEEIFEKHADV
 IFEEARNRLYVVKAALLCFLDNQRGRE

SEQ ID NO:24

ACE 941 Cj0998c

>AnrP979073 (NC_002163) putative periplasmic protein [C. jejuni]
 MKKILVSVLSSCLLASALSASVSKEDSLKISFEGYKTKDMIGTKGEFKNVEYKFSKNIKDLASYLKGAKA
 TIKPSNAFMGEGNIDIITNNITKVFFPALLGDTDIKVVFQDVAGENKGVISAKITMDKSTIVPLTYTIK
 DNKFEAKGQQLDLHTFKNGSKALKALSDVAAGHGGISWPVLDISFNADLAE

SEQ ID NO:25

ACE 961 Cj1018c

>AnrP257863 (NC_002163) branched-chain amino-acid ABC transport system periplasmic binding protein [C. jejuni]
 MKKSLILASILSLSLAAEVKIGVVLPLSGATAAYGQSALEGIKLANSQMSALSGDKVSLAIIDTKGDK
 LESSSGANRLVSQDKVIGLIGEMVTANTLQVMRVAEDNPKIPLIAPAATGDRLLDKKIYSSRVCMDSFQG
 SSLAKYVFSKLNYSKSAVIVDQSTDYSLGLAKAFEQYKSNGGQILRILRVNSGDKDFRAIVAQVKSLSNP
 EFIGFLPLYSEASLFARQSKLAGLNIPMGSADGVADQTFIGSLAGDASEGYIFTDSFDANNPTTKLSKEFI
 SVYEKAKGTKEVPNFSAMGADAYFVMLNAMNACVENLTSKCVNEKIHQTKNYQGVSGVISIDQTGNATRS
 VVVKEIKNQKQNYKDIINP

SEQ ID NO:26

ACE 962 Cj1019c

>AnrP326257 (NC_002163) branched-chain amino-acid ABC transport system periplasmic binding protein [C. jejuni]
 MKKLTTLTSLVLMVNCLYAKDINIGVVLPLTGTVAAYGQDVNGIELANKLQPKLSNGDVVIKLITIDTKG
 DKLETSGVNRLIATDKVLGIIGEATTPTNTIQAISIAEKKIPLIAPVASGDKLIDKKKYASRVCFKDSF
 QGDKFATVYVSKDLGLKNAVIIIDQSNVYSLGLARAFENSFKNNNGKIIKKLVINSGDKDFRAVVSQLKSL
 NPDFVYMPYHPEAALIARQARQIGFDKLLVAGDGVNNQTFIDLGGSAVNGVIFTDSFDYNSPSTQLGKD
 FVAAYEKVKGTKEPAFSAMGADAYFVMLNAMNACVDNLSECINSKIHQTKDFQAVGGVISIDESGNAI
 RSVVIKEIQNQKQNYKTIINP

SEQ ID NO:27

ACE 984 Cj1041c

>AnrP198268 (NC_002163) putative periplasmic ATP/GTP-binding protein [C. jejuni]

MKKYVLSLALLGSSLGASELKQYEFDGFKSPESIFVVDKNYVYVSVNVGEKLEPLAKDNDGFISKLDKNGKV
 LEYKFLTHLNAPKGMMEIGKTLYVVDIDVLRGFDLKTKEIFNLPIKGAIFLNDIEKLDDNTLLVSDTGT
 GLILKVDLTKTQYDELLKLDLAKFGGPNGLYLDRKKHKLFIAGYHPDGVSggvVMayDLNTKELSIIKNE
 KESYDGIVPYKDGLLVSWSGNLNGYIYNLDNVKSVKLEPLIMKGPADIEGNILWIPKMVEGKIFKVE
 LNK

SEQ ID NO:28

ACE 1094 Cj1153

>AnrP385049 (NC_002163) putative periplasmic cytochrome C [C. jejuni]
 MKKLLVVSALA CLGVSAFAADGATLFKKCAVCHGANADKVYLNKPALKTLSSAERLQYMKEYSEGKRNA
 YGQGAIMKLNLKGLTEEDFKIAEHIAETLK

SEQ ID NO:29

ACE 1155 Cj1214c

>AnrP470247 (NC_002163) hypothetical protein Cj1214c [C. jejuni]
 MFKTIVCFALNLSSLFAVGFDLKPPIKSELVKVDDIYGYIKDSDDIKLYSSGVVVQHFNSQSIIARASVI
 DKKNGLAKLEFSVFSALKQDALPLPNVLPKVGDEVVLFYDRGLVIAPDEQTYNELVREFPQIYFTHID
 IFGAQLIRTATLSPKRSDFRQFCDDNAVGLVVALENHAEVVDQCDFNKLYEVPISKPTSVQVPFYSRIG
 GYKSNFFDFNSQEIGNYYRYYDALINLPKVQ

FIG. 4

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SEQ ID NO:30

ACE 1227 Cj1287c

>AnrP530915 (NC_002163) malate oxidoreductase [C. jejuni]
 MNLKEEALKYHLLGGKIDIVPSKPMATSYDSLALAYSPGVAEPCLIEAKDNELAYTYTNKANLVAIVSDGSA
 VLGLGNIGAQASKPVMEGKACLFKKFANVNAVDIEINVHSIEEIVNFCKALAPTVGGINLEDIAAPKCFE
 IEAALQDLGIIPVMHDDQHGTAAIISTAGLMNAMEISGKFKDICKVVSGAGAAGIASAKMYRNLGVENIIL
 VDSKGVIISKDRNDLTPQKLEFAVDSKEKTLKEVLKGADVFLGLSAPKILDDEMVLMSMAKDPVIFALANPI
 PEVMPEDVARLRKDAIVGTGRSDYPNQINNVLGFFIFRGALDVRAASKITENMKVAAAKALADLAKLPVS
 DAVKKAYNLSTLEFGRDYVIPKPFDERVKAAVSTAVAAAACKDGVAKVKNFDEKAYFESLK

SEQ ID NO:31

ACE 1320 Cj1380

>AnrP108083 (NC_002163) putative periplasmic protein [C. jejuni]
 MKKLSLILVCSASLFAASNSEISDFYSKSISKAQFPNATSVSVSRQKVGNTRFESVIVSVELNGQKQENIL
 FTKDSLITPDIDLKTGISYAQEYEMKKFQEARENFTKNAKAVAQKETMIALGDKNKPAIYVFSDPECP
 YCREHLAQIDDELKNYQVNYILTPTVHGKSAFEKSALITYKEAKKAKNDKEKIAILNKYYDANIKNYPKVSD
 AELKEVFSLYEKYRSLGLSATPTIIK

SEQ ID NO:32

ACE 1431 Cj1496c

>AnrP947055 (NC_002163) putative periplasmic protein [C. jejuni]
 MIKKFILLVFISSVVFGAEQDCEQYFEARKAQIELQTRFDEARQSLEAYKASFEALQKERLENLEKKEA
 EVNATLAKIEELKLENARLVEEQQKILNSINDKTQGRVKEIYSQMKDAAIADVLSQMDAEDASKIMLSLE
 SRKISGVLSKMDPKKASELTLLLKNDNNNASN

SEQ ID NO:33

ACE 1449 Cj1516

>AnrP407504 (NC_002163) putative periplasmic oxidoreductase [C. jejuni]
 MNRRNFLKFNLNTLASMGVAYANPMHDHMHSMHKNHSINHLDLTSFINFAPKPNLKLDPKQFPQGEILKAL
 PLLKNESEKKNIFRATLEIENHIELIKGKTLFYTYNGLVPAPKIEVPEGDKLEILVKNKLKEATTIHW
 HGVPVPPDQDGSPHDPIAGEERIYRFEIPQDSAGTYWYHPHPYTAKQVFMGLAGAFVIKAKKDALS
 LKEKDLMSIDRLDENAQIPNNNLNDWLNGREGEFVFLINGQFKPKIQLATNERIRIYNATAARYLNRIQ
 GAKFILVGTDGGLIEKAIYKEELFLSPASRVEVLIDAPKDGNFKLESAYYDRDKMMVKEESTNLFLANIN
 LKKEKLELPKNLKIFKPLEEPKEFKEIIMSEDHMQMHHGMGKSENELKIALASMFLINGKSYDLKRIDLS
 SKLGVVEDWIVINKSHMDHPFHNGTQFELISSKLNQVQKAERFRDTINVRPNEELRLKMQDFKGL
 RMYHCHILEHEDLGMGNLEVKE

SEQ ID NO:34

ACE 1469 Cj1540

>AnrP818860 (NC_002163) putative periplasmic protein [C. jejuni]
 MKKIISLALALALSASAELKMMTTSTDNTGLLDALKPLYEKESENGLKWWAVGTGAALKMGEDCNADV
 LFVHSPKAKEEFMKGFGVDRTPMYNDFIIAADKSLASKFKGKLNKESLELIKNEKLTFSRGDKSGTD
 NKEKSLWKNLGGVPEKQSWYQQSGQGMLASIKIAEEKKVILTDRTGYIYKEANEKGKPVLVIVNEGDDS
 LKNFYSVIATNPKHCKNVNYTEASKFIKWTSKTLNFIADFKLNNKPLFVIDAKTRKD

SEQ ID NO:35

ACE 1584 Cj1659

>AnrP294550 (NC_002163) periplasmic protein p19 [C. jejuni]
 MIKKVLSVAAAIVISTNLFAGEVPIDGPKELNGMEIAAVYLQPIEMEPRGIDLAASLADIHAL
 KNNPNGFPEGFWMPYLTIAYELKMTDTGAIKRGTLMPVADDGPHYGGANIAMEKDKGGFVGVNYELTFY
 ISNPEKQGFGRHVDEETGVGKWFEPFKVDYKFKYTGTGPK

SEQ ID NO:36

ACE 1569 Cj1643

>AnrP407676 (NC_002163) putative periplasmic protein [C. jejuni]
 MKKILITCMLFTSFGIERPKFEDFLAGYERNKASMLNYEGMPAALSENLLAVLKQPNAKLKYVKYD
 FLNLYLVRTDFSLIPTPMGDEEKLTRNDWVGIWDPNPKYIGHIKYLAQNIDEKDQDFNSKIGLLGTPCC
 EMMGIALNNSSFIGNRYLKHFMKYNDVYWDGIVGDFVVRRENKIYVNNVRKNPQFLINDQVISVDGLPAND
 LRKLNEKILFADRGSTLYFQVLRDMDLNISTEVFAKDLISKFKNLPDSKPKPKITNFTSNLGLTVNASLVV
 TKIDPKSKVSNAGFMVGDKILRVNNIILNNFKELQNILSAGNDFSILIERKSTKLPLSNFNNELGGNANS
 GGDGKFQFFIRLTK

SEQ ID NO:37

ACE 96 Cj0105

>AnrP758295 (NC_002163) ATP synthase F1 sector alpha subunit [C. jejuni]
 MKFKADEISSLKERIENFDLNLEIEETGKIIISVADGVAKVYGLKNIMAGEMVEFENGDKGMALNLEESS
 VGIVILGKGEGLKEGASVKRLLKVPVGEALIGRVVNALGEPIDAKGVINANEYRFVEEKAKGIMARK

FIG. 4

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SVHEPLHTGIKAIDALVPIGRGQRELIIGDRQTGKTTVAVDTIISQRGQGVICIVYVAIGQKQSTVAQVVK
 RLEEHGAMEYTIVVAGASDPAALOYLAPYTGVTMGEFFRDNAKHALIVYDDLSKHAVAYREMSLILRRP
 PGREAYPGDVFYLHSRLLERASKLNDELGAGSLTALPIETQAGDVSAYIPTNVISITDGQIFLETDLFN
 SGIRPAINVGLSVSRVGGAAQIKATKQVSGTLRLDAQYRELQAFQFASDLDEASRKQLERGQRMVELL
 KQPPYSPLSVEKQVVLIFAGTKGFLDDIAVSRIKEFEDGIYPFIEAKHPDIFEQIRSKKALDSLEEKLA
 KAINEFKANHL

SEQ ID NO:38

ACE 165 Cj0175

>AnrP550554 (NC_002163) putative iron-uptake ABC transport system
 periplasmic iron
 MKKIFFMFMLTAWSFLGASELNTIYSARHYNADFEIICKFEEKTGIKVNHQTQAKASELIKRLSLEGSNSPAD
 IFITADISNLTEAKNLGLLSPVSKYLEEPIPAHLRDKDKEWFAITKRARIIAYNKNTNIDISKMKNYED
 LAKAEFKGEIVMRSATAPYSKTLASIIANDGNKEAKAWAKGVLENLATNPKGGRDQARQVFAGEAKFA
 VMNTYYIGLLKNSKNPKDVEVGNGLIIFPNQDNRGTHINISGIAINTKSSKNQDAAKKFMEFMLSPEIQL
 ILTDSNYEFFPIRNDVVELSQTVKDFGTFKEDQIPVSKIAENIKEAVKIYDEVGFR

SEQ ID NO:39

ACE 257 Cj0283c

>AnrP602342 (NC_002163) chemotaxis protein [C. jejuni]
 MSNEKLEQILQKQQTQMAGPDVDQREDDIIQLVGFVVVGDEEYAIPILNQELIIPKIEYTRVPSVPDYVLG
 VFNMRGNVMPPLIDLAQRFHLSKMTQPTRYIVLRGETNGTGVGGNAGFVIDRLTEAIKIHRNRIDPPPE
 TLVKDKGMIVGIGKRDENILTILKVEALLKREF

SEQ ID NO:40

ACE 277 Cj0303c

>AnrP311344 (NC_002163) putative molybdate-binding lipoprotein
 [C. jejuni]
 MKKFVVFPGILLFVSCCLNAQNLISIVFVASSASKAMSEVKDEFLKTHPEDKIELVFGASGKYELLKQGREF
 DLFFSADTKYAKAIYDDKNALIKPKVYVLGVLAlysLDENLLQGGVENLKEKANKITHLSIANPKVAPYG
 VAAKEVLENLGLNELLKDKIVLGENISVFPVLHVDSKNSDIAIVAYSLVSSINHPKGKAVIIDAKYFSPLE
 QSYVITKYAKDKKLAFEFNEFIGSSKAKEIFKKYGFSTP

SEQ ID NO:41

ACE 308 Cj0334

>AnrP505685 (NC_002163) alkyl hydroperoxide reductase [C. jejuni]
 MIVTKKALDFTAPAVLGNNEIVQDFNLYKNIGPKGAVVFFYPKDFTVCPSEIIAFDKRYQEFKNRGIEV
 IGIISGDNEFSHFAWKNTPVNQGGIGQVVFPLVADLTQKQIARNFDVLYAEAVALRGSFLLDADGTVRHAVV
 NDLPPLGRNIDEMLRMVDTMLFTNEHGEVCPAGWNKGDEGMKANPKGVAEYLGKNEAKL

SEQ ID NO:42

ACE 388 Cj0415

>AnrP72219 (NC_002163) putative oxidoreductase subunit [C. jejuni]
 MAEVLLKVDVVTVGAGWTGGIVAAELTKAGLNVLSLERGHMQSTENFNYIHDEWRGYINYGLMQDCSKDT
 VTFRHDPSGLALPYRKMGFLLGNVNGGAGVHWNGWTFRFMPYDFEIQTLSKQRYGNKLGNDYTLQDWGV
 TYKDMEPYYDRFEKTCGVSGEPNPLAEMGAFRSSPYPQEPLENTKMLKRFESAACKSSNLHTYRLPASN
 KGGYTNPDGQD LAPCQC YCAYCERFGCEYAKASPLNTVIPKAMSTGKYTIRTYSNVTQILKKDGKVTV
 FVDTRTMKEYIQPADIVVLT SYMFNNAKLLMVS NIGEQYDPKTKGKTLGRNYCYQMNMGTTAFFDEQFNT
 FMGSGALGTTSDDFNGDNFDHSKEKFLHGAMIYSVQLGTRPIQOSA PLPAGAPT WGA EFKKALN
 YNFTRAI TVGGQGASLPHKNNYLSLDPTYKDAFGMPLRL TYNFTDQDRA LHKFITDKTAEVAKRMQGV
 KSIKKGAY LKDYSVVPYQSTHNTGGTTMGADRETSVNVNTYLOHWDADNLFVVGAGNFQHN
 SGINPTDTV GALAYRCAE GILKYHKSGKSLA

SEQ ID NO:43

ACE 393 Cj0420

>AnrP490750 (NC_002163) putative periplasmic protein [C. jejuni]
 MKVLLSSLVAVSLLSTGLFAKEYTLDKAHTDVGFKIKHLQISNVKGNFKDYSAVIDFDPASAEFKKLDX₁
 TIKIASVNTENQTRDNHLQODDDFKAKKYPDMFTMKKYEKIDNEKGKMTGTLTIAGVSKDIVLDX₂EIGG
 VAKGKDGEKIGFSLNGKIKRSDFKFAX, STSTITLSDDINLNIEVEANEK

wherein X₁ is V or A, and X₂ is A or T, and X₃ is T or A

SEQ ID NO:44

ACE 711 Cj0759

>AnrP586832 (NC_002163) heat shock protein dnaK [C. jejuni]
 MSKVIGIDLGTTNSCVAVYERGESKVIPNKEGKNTTPSVVAFTDKGEVLVGDASKRQAVTNPEKTIYSIK
 RIMGLMINEDA AKEAKNRLPYHITERNGAC AIEIAGKIYTPQEISAKVLMKLKEDAE AFLGESVTD
 AVIT

FIG. 4

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VPAYFNDQAQRKATKEAGTIAGLNVLRIINEPTSAALAYGLDKKDSEKIVVYDLGGGTFDVTVLETGDNVV
 EVLATGGNAFLGGDDFDNKLIDFLANEFKDETGDILKNDVMALQLKEAAENAKKELSSANETEINLPFI
 TADASGPKHLVKKLTRAKFEGMIDSVAETITKINEVVDAGLKKDEIKEIVMVGGSTRVPLVQEEVKKA
 FNKDLNKSVNPDEVVAIGAAIQAVIKGDVKDVLLDVTPLSLGIETLGGVMTKIEKGTTIPTKKEQVF
 STAEDNQSAVTINVLGGERFSRDNKSLGNFNLEGIPPAAPRGMPQIEVTFDIDANGILTSAKDKATGKA
 QEIKITGSSGLSEEEINNMVKDAELHKEEDKCRKEAVDARNAADSLAHQVEKSLSELGEKVAADKENIQ
 KALDDLRETLKNQNASKEEEIESKMKALSEVSHKLAENMYKDEPTANDKKKDDVIDAEVE

SEQ ID NO:45

ACE 723 Cj0771c

>AnrP524051 (NC_002163) putative periplasmic protein [C. jejuni]
 MNLFKIIIILACILNLSSLFAQONITIGATPNPGSLLIELMKDDFKNKGYELKIVEFSDYILPNRALEEKEL
 DANLYQHKPFLEEYNLKKGSNLIAATTPLVIAPIVGVYSSKKIKNLENLKEGARVAIPNDATNESRALEEK
 AKLIELNKNTLKTPLDINKPKKLKFIELDKAAQLPRALEDDVDIAIINSNFALGAGLNPSKDTIFREDKNS
 PYVNYVVVRSEGKNSEKTKVIDEILRSDFKAIINEHYKDILIPAF

SEQ ID NO:46

ACE 724 Cj0772c

>AnrP579672 (NC_002163) putative periplasmic protein [C. jejuni]
 MKIKSLFIAISILTLSNANALETITVAATPVPHAEILEQVKPDLEQGYKLEIKEFTDYVLPNLAVDNGE
 ADANFFQHTPYLEEFNKNKGTLIKVAAIIEPMAVYSKKYKSLDDIKEGVKIAIPNDPTNESRALDI
 KKGLVFKFKDALKTPLDIIDNPKKIKFVELKPAQLPRALEDDVDIAIINSNFALGAGLNPSKDTIFREDKNS
 SPYANILVVVRVGHENDPKIKALIQALQSDKIKQFIIKEYNGSVPFAF

SEQ ID NO:47

ACE 819 Cj0872

>AnrP694298 (NC_002163) putative protein disulphide isomerase
 [C. jejuni]
 MRNFFCKFVLALVFYSSFALANNSFITLNPSLPSSENSVIEAFSYKCIHCYNNHHKFGTLEKLREAFPNLH
 FKLYPVSIMNGEFSKEMNELFTFAQYKDEQNGKDASYSDSLSHKLADVYFVSYFLNKQRNFSNLDEFYDI
 GLKAMNVNKNEVLNFIINTPKAKEILSEFQRANDIACYGTGPAFVVNGKYQINPSAINSMQDLEDLVKKLS
 NMK

SEQ ID NO:48

ACE 1060 Cj1118c

>AnrP515430 (NC_002163) chemotaxis regulatory protein [C. jejuni]
 MKLUVVDDSSSTMRIIKNTLTRLGHDDVLEAEHGVEAWLLTKNEDVKVLITDWNMPEMNGLELVKKVRA
 EKKYEDMPIIMVTTEGGKAEVITALKAGVNYYIVKPFTPQLKEKLEDVLGTSAGEAAE

SEQ ID NO:49

ACE 1169 Cj1228c

>AnrP679791 (NC_002163) serine protease (protease DO) [C. jejuni]
 MKKIFLSLSLASALFAASINFNESTATANRVNPAGNAVLSYHDSIKDAKKSVNI
 STSKTITRANRPSP
 LDDFFNDPYFKQFFDFDSQRKGKNDKEVSSLGSGVIISKDGYIVTNNHVVDADTITVNLPGSDIEYK
 AKLIGKDPKTDLAVIKIEANNLSAITFTNSDDLMEGDVVFALGNPGVGFVTS
 SGIIISALNKN
 DNIIGLNQY
 ENFIQTDASINPGNSGGALVDSRGYLVGINSA
 LSRG
 GGNNGIGFA
 IPSNMVK
 DIAKK
 LIEKGK
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SEQ ID NO:50

ACE 1510 Cj1584c

>AnrP355324 (NC_002163) putative peptide ABC-transport system
 periplasmic peptide- [C. jejuni]
 MLRWFVLLFLNLEAKIPIKPDLLTIIAVENETIARINPAYSEDHD
 AVINLVFSGLTRFDENMSLKPD
 LAKS
 WD
 ISKDG
 LVYDIFLR
 DDLWHDGV
 KFSADDV
 KFSI
 LEAF
 KNPK
 N
 SS
 IYV
 NF
 ED
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SEQ ID NO:51

ACE 1543 Cj1617

>AnrP111949 (NC_002163) putative haemin uptake system periplasmic
 haemin-binding protein [C. jejuni]

FIG. 4

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MKKILIIIMSLFLIALNAKERLVVLDPASIELTFLMLKAEDQIVGIATLQHSNIYPKDQTSKLTSGTFSNP
 SLEKIVALKPSLVLVILSSYSLNLEEGLKNFGIKSINLKAERLEDITKNITLGGQITKKEKEAELLKQEFNQ
 KFKKLSDKPLNKSAIYLSSNPLMAFDNSLIADILRLIGIKNLSPSQISRPVISAEYILKQNPDILIL
 GINAKNNLLDTNALLKNTKAVKTGSIYFNKDTPILLRLSPKIIDRIQEFTKLENNNF

Fragments of AnrP630851 (Cj0092)

SEQ ID NO:52: TQQDINTQNEMSDASTKDITPKSIEDFFEEFAD
 SEQ ID NO:53: GITKDGKTFYTGKSTVAVNDTDPQF
 SEQ ID NO:54: RIATSKIQNYEADNSTNAKEFDELPKGDKVDQILNK
 SEQ ID NO:55: QLDKALKDLGIDTNLSLEDRKKTLLKQEFLNKTMTN
 SEQ ID NO:56: TIVTQRGRGEYDVGVVAVISNKTRQLAKD
 SEQ ID NO:57: AISEYLPKDTKGFLNEYGIRLVYDEN
 SEQ ID NO:58: DPSNAKKTNILEDRAKET
 SEQ ID NO:59: LSLKDERTTGDTYEEII
 SEQ ID NO:60: VNDSSSTQEQTQNITN
 SEQ ID NO:61: TLKKWSYTSENG
 SEQ ID NO:62: YSYENLANTNEALNSKSNAKNEAKKSSSIQRS

Fragments of AnrP57234 (Cj0143c)

SEQ ID NO:63: IFYTFTQAKNLEQEQQNTSSNLVSVS
 SEQ ID NO:64: LPPNSNEHNFEFKPSTMKKLEKSDIYF
 SEQ ID NO:65: LEFEKVFTDKFKQNFPKLQVINMQ
 SEQ ID NO:66: IQTHDTHEHSHEHHHEHGHFDPTHWL
 SEQ ID NO:67: DTЛИQKPQNEONLYKENLDK
 SEQ ID NO:68: SKLEKLKNRE
 SEQ ID NO:69: YFAKRYNL
 SEQ ID NO:70: GKEPKSKDQK
 SEQ ID NO:71: LMKDKNLK
 SEQ ID NO:72: QNGFPENAATLAKEDAKIYK
 SEQ ID NO:73: DHLSYDWEWELLKTADA

Fragments of AnrP490750 (Cj0420)

SEQ ID NO:74: FAKEYTLDKAHTDVGFKIKHLQI
 SEQ ID NO:75: VKGNFKDYSAVIDFDPASAEFKKLDVTI
 SEQ ID NO:76: SVNTENQTRDNHLQQDDFF
 SEQ ID NO:77: DFFKAKKYPDMDTFTM
 SEQ ID NO:78: TFTMKKYEKIDNEKGKMT
 SEQ ID NO:79: GVAKGKDGEKIGF
 SEQ ID NO:80: LNGKIKRSDFKFATS
 SEQ ID NO:81: EVEANEK

Fragments of AnrP684299 (Cj0715)

SEQ ID NO:82: LSATEYQLSTHV
 SEQ ID NO:83: ITSGQPAPKVVKVELYK
 SEQ ID NO:84: LEANQQWKVSEEFTEENGRIG
 SEQ ID NO:85: LLPYEKAENRAFG
 SEQ ID NO:86: KFFTKDYYTSHKINTF
 SEQ ID NO:87: SFELSKDQKHYPVPI
 SEQ ID NO:88: FGYSTYRGS

Fragments of AnrP579672 (Cj0772c)

SEQ ID NO:89: ILEQVKPDLKEQGYKLEIKEFTDY
 SEQ ID NO:90: GEADANFFQHTPYLEEPNKNKGT
 SEQ ID NO:91: AVYSKKYKSLDDIKE
 SEQ ID NO:92: IPNDPTNESRAL
 SEQ ID NO:93: KGLVKFKDKALKTPLDIIDNPKKIKFVELKPAQLPRALN
 SEQ ID NO:94: ANLNPAKDSVFIEDKESPYAN
 SEQ ID NO:95: GHENDPKIKALIQALQSDKIKQFIIEKYN

Fragments of AnrP257863 (Cj1018c)

SEQ ID NO:96: NGDKVSLAIIDTKGDKLESSGANRLVSQLK
 SEQ ID NO:97: VAEDNKIPLIAPAATGDRLLDKKIYSSRVC
 SEQ ID NO:98: YVFSKLNKSAVIVVDQSTDYSLGLAKAFEKQYKSNGGQ
 SEQ ID NO:99: NSGDKDFRAIAVQVKSINPEFIFLPLYSEASLFARQSKLA
 SEQ ID NO:100: GYIFTDSFDANNPTTKLSKEFISVYEKAGTKEVPNFSAMG

FIG. 4**19/20**

SEQ ID NO:101: VNEKIHQTKNYQGVS

SEQ ID NO:102: QTGNATRSVVVKEIKNQKQNYKDIIN

Fragments of AnrP108083 (Cj1380)

SEQ ID NO:103: ELNGQKQENILFTK
SEQ ID NO:104: SYAQEYEMKKFQEARENFTKNAKAVAQKETM
SEQ ID NO:105: VIALGDKNKPAI
SEQ ID NO:106: HLAQIDDELKNKYQV
SEQ ID NO:107: YKEAKKAKNDKEKIAI
SEQ ID NO:108: LNKYYDANIKNYPKVSDAELKEVFSLYEKYRSL

Fragments of AnrP407676 (Cj1643)

SEQ ID NO:109: GIERPKFEDFLAGYERNKASMLN
SEQ ID NO:110: LKQPNAKLNKVVKYDPFL
SEQ ID NO:111: PTPMGDEEKLTRNDWVGWIWDPNPKPYI
SEQ ID NO:112: AQNIDEKDQLDFNSK
SEQ ID NO:113: GNRYLKHFMKYNDVY
SEQ ID NO:114: VVRENKIVNNVRKNPQFL
SEQ ID NO:115: PANDLRKLNEK
SEQ ID NO:116: DRGSTLYFQVLRDNMDLN
SEQ ID NO:117: AKDLSKFNLPDSPKPKI
SEQ ID NO:118: TKIDPKSKVSNAG
SEQ ID NO:119: LIERKSTKLPLSBNFN

SEQ ID NO:120:

ACE 393, ML53 (0:19)DNA

ATGAAAAAAAGTTTATTGAGTTCAATTGGTTGGGTGTC TTGTTAACAGCACAGGTTGTTGCTAAAGAAATATACTTTAGA
TAAAGCACATACAGATGTAGGTTAAAATCAAACATTACAAATTAGCAATGTAAAAGGAAATTCAAAGATTATCTG
CGGTGATTGATTGGACCCCTGCAGTGCTGAATTAAAAGCTTGATGCAACTATAAAATCGCATCTGTAATACAGAA
AATCAAACAAGAGATAATCACTTACAGCAAGATGATTTCAAAGCAAAAAAATATCCTGATATGACTTTACAATGAA
AAAATATGAAAAAAATCGATAATGAAAAAGTAAAATGACAGGAACTTTAACTATAGCTGGAGTTCTAAAGATATCGTT
TAGATACTGAAATCGCGGTGTAAGCTAAAGGCAAAGATGGAAAAGAAAAAATAGGCTTTCTTAAATGGAAAATCAAA
CGCTCTGATTAAATTGCAAGTACTTCAACTATTACTTAAAGTGTATTAATTAAATATCGAAGTTGAAGC
GAACGAAAAA

SEQ ID NO:121: DIVLDTEIGGVAK

SEQ ID NO:122: FAASTSTTLSDDINLNIEVEANEK

SEQ ID NO:123: LDATIK

20/20

Table 1